Human TCR V α -1.5 (V α -8.2) coding sequence

AACCAGAGCC CAGTCGGTGA CCCAGCTTGA CAGCCACGTC TCTGTCTCTG ATGCTCCTGC TGCTCGTCCC AGTGCTCGAG GTGATTTTTA CTCTGGGAGG SAAGTACACA TCAGCGGCCA CCCTGGTTAA AGGCATCAAC GGTTTTGAGG AGGAGGTGCT GACGGACTCA CCTTTGGCAA AGGGACTCAT CTAATCATCC FGCTAGACAT GAGGTCTATG GACTTCAAGA GCAACAGTGC TGTGGCCTGG CATATGAGCG ACGCGGCTGA GTACTTCTGT GTTGTGAGTC CTTTTTCAGG AGCCCTATAT CCAGAACCCT GACCCTGCCG TGTACCAGCT GAGAGACTCT STGAATTTAA GAAGAGTGAA ACCTCCTTCC ACCTGACGAA ACCCTCAGCC AAGGAACCCC GGTGCTGCTG AGGTGCAACT ACTCATCTTC TTATTCACCA CCAGAAGAC ACCTTCTTCC CCAGCCCAGA AAGTTCCTGT GATGTCAAGC CAGTGATTG GGTTCCGAAT CCTCCTCCTG AAAGTGGCCG GGTTTAATCT GGTCGAGAA AAGCTTTGAA ACAGATACGA ACCTAAAACTT TCAAAACCTG ICTCTCTTCT GGTATGTGCA ACACCCCAAC AAAGGACTCC AGCTTCTCCT AAATGTGTCA CAAAGTAAGG ATTCTGATGT GTATATCACA GACAAACTG AAATCCAGTG ACAAGTCTGT CTGCCTATTC ACCGATTTTG ATTCTCAAAC AGCAACAAAT CTGACTTTGC ATGTGCAAAC GCCTTCAACA ACAGCATTA⁻ SCTCATGACG CTGCGGCTGT GGTCCAGCTG A

Rgure 2

Human TCR V α -1.5 (V α -8.2) protein sequence

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PVLLRCNYSSSYSPSLFWYVQHPNKGLQLLLKYT CDR1

MLLLLVPVLEVIFTLGGTRAQSVTQLDSHVSVSEGT

CDR2

SAATLVKGINGFEAEFKKSETSFHLTKPSAHMSDA

FR3

AEYFCVVSPFSGGGADGLT CDR3

TFFPSPESSCDVKLVEKSFETDTNLNFQNLSVIGFRIL FGKGTH LIIOPYIONP DPAVYQLRDSKSSDKSVCLF TDFDSQTNVS QSKDSDVYIT DKTVLDMRSM DFKSNSAVAWSNKSDFACAN AFNNSIIPED LL KVAGFNLLMT LRLWSS

Human TCR Vβ-2.1 (Vβ-20.1) coding sequence

GGAGACCCAG TACTTCGGGC CAGGCACGCG GCTCCTGGTG CTCGAGGACC CTACCCCGAC CACGTGGAGC TGAGCTGGTG GGTGAATGGG AAGGAGGTGC ACAGTGGGGT CAGCACAGAC CCGCAGCCCC TCAAGGAGCA GCCCGCCCTC ATCGTCAGCG CCGAGGCCTG GGGTAGAGCA GACTGTGGCT TCACCTCCGA GAAGACAGCA GCTTCTACAT CTGCAGTGCT AGAGATGGGG GGGAGGGTTC TCTCGGAGAA TGACGAGTGG ACCCAGGATA GGGCCAAACC TGTCACCCAG FAGGGAAGGC CACCTTGTAT GCCGTGCTGG TCAGTGCCCT CGTGCTGATG ATGCTGCTGC TTCTGCTGCT TCTGGGGCCA GGCTCCGGGC TTGGTGCTGT CTGGCAGAAC CCCCGCAACC ACTTCCGCTG TCAAGTCCAG TTCTACGGGC FGAAAAACGT GTTCCCACCC GAGGTCGCTG TGTTTGAGCC ATCAGAAGCA GAGATCTCCC ACACCCAAAA GGCCACACTG GTGTGCCTGG CCACAGGCTT CTCCAAGGCC ACATACGAGC AAGGCGTCGA GAAGGACAAG TTTCTCATCA CGTCAGTTCC CGAAACAGAG TCTCATGCTG ATGGCAACTT CCAATGAGGG AATGACTCCA GATACTGCCT GAGCAGCCGC CTGAGGGTCT CGGCCACCTI CGTCTCTCAA CATCCGAGCT GGGTTATCTG TAAGAGTGGA ACCTCTGTGA GTCTTACCAG CAAGGGGTCC TGTCTGCCAC CATCCTCAT GAGATCTTGC ACCATGCAAG CCTGACCTTG TCCACTCTGA CAGTGACCAG TGCCCATCCT AGATCGAGTG CCGTTCCCTG GACTTTCAGG CCACAACTAT GTTTTGGTAT GCCATGGTCA AGAGAAAGGA TTCCAGAGGC TAG

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Figure 4

Human TCR V β -2.1 (V β -20.1) protein sequence

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MLLLLLLGPGSGLGAVVSQHPSWVICKSGTSVKIECR

SLDFOATTMFWYRQFPKQSLMLMATSNEGSKATYEQ CDR1

GVEKDKFLINHASLTLSTLTVTSAHPEDSSFYICSARD FR3

CDR3 GGEG

KATLVCLATGFYPDHVELSWWVNGKEVHSGVSTDPQPL GLSENDEWTQDRAKPVTQIVSAEAWGRADCGFTSESYQ **QGVLSATILYEILLGKATLYAVLVSALVLMAMVKRKDS** SETQYFGPGTRLLVLEDLKNVFPPEVAVFEPSEAEISHTO KEQPALNDSRYCLSSRLRVSATFWQNPRNHFRCQVQFY constant

Human TCR V α -1.5 (V α -8.2) protein sequence

Figure 5

MILLLVPVLEVIFTLGGTRAQSVTQLDSHVSVSEGT

PVLLRCNYSSSYSPSLFWYVQHPNKGLQLLLKYT CDR1

SAATLVKGINGFEAEFKKSETSFHLTKPSAHMSDA

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AEYFCVVSPFSGGGADGLTFGKGTH LIIQP 145 CDR3 Va8.2

TFFPSPESSCDVKLVEKSFETDTNLNFQNLSVIGFRIL YIQNP DPAVYQLRDSKSSDKSVCLF TDFDSQTNVS **DFKSNSAVAWSNKSDFACAN AFNNSIIPED ƏSKDSDVYIT DKTVLDMRSM** LL KVAGFNLLMT LRLWSS constant

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Human TCR VB-2.1 (VB-20.1) protein sequence

figure 6

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MILLILLIGPGSGLGAVVSQHPSWVICKSGTSVKTECR

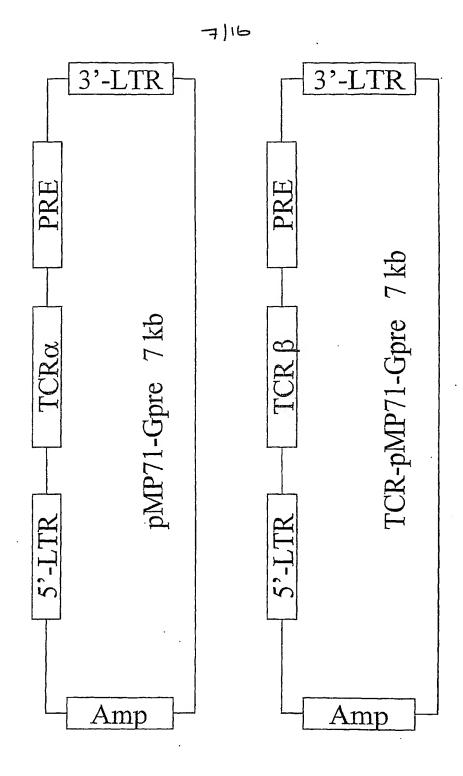
SLDFQATTMFWYRQFPKQSLMLMATSNEGSKATYEQ CDR2 FR2

GVEKDIKFLINHASLTLSTLTVTSAHPEDSSFYICSARD

CDR3
GGEGSETQYFGPGTRLLVL

RLRVSATFWQNPRNHFRCQVQFYGLSENDEWTQDRAKP EDLKNVFPPEVAVFEPSEAEISHTOKATLVCLATGFYPDH VELSWWVNGKEVHSGVSTDPQPLKEQPALNDSRYCLSS VTQIVSAEAWGRADCGFTSESYQQGVLSATILYEILLGK ATLYAVLVSALVLMAMVKRKDSRG Constant 2

TCR-retroviral constructs



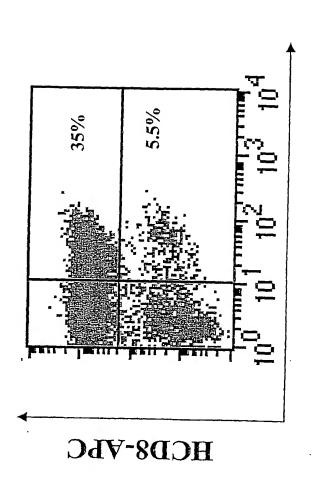
HCD8-VbC

HVB2.1-FITC

Figure

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Increase of CD8⁺-Vb2.1⁺ T Cells after antigen stimulation



HVB2.1-FITC

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TCR specific for pWT126 transduced PBMC

CD8

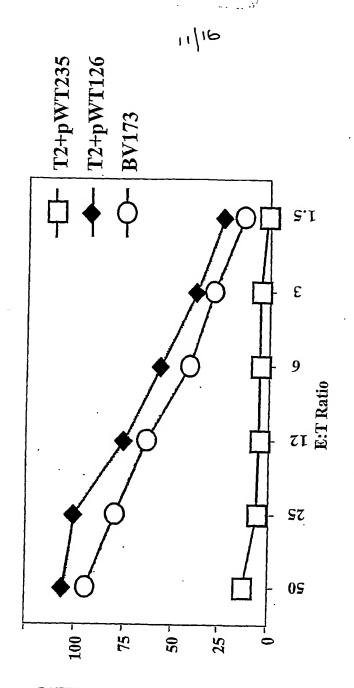
CD8

Tetramer

Tetramer

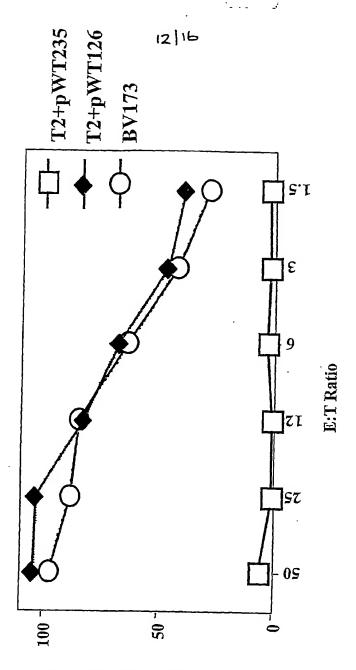
Figure (0

TCR transduced bulk T cells show pWT126-specific killing activity



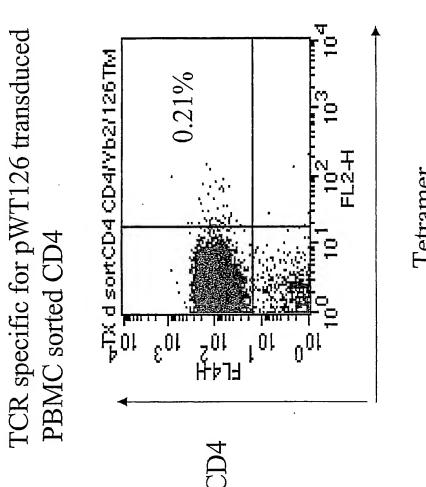
% SLECILIC KITTING

TCR transduced CD8+ T cells show pWT126-specific killing activit



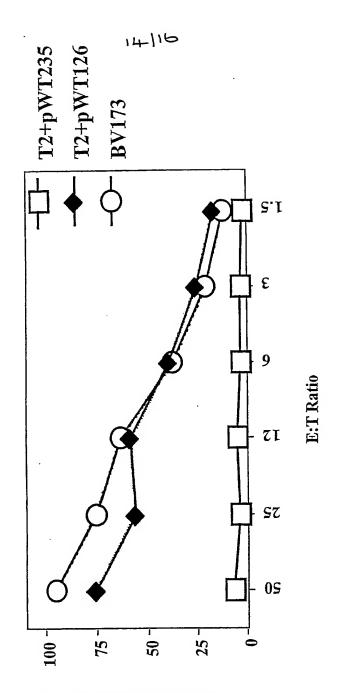
% SEECIEIC KITTING

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Tetramer

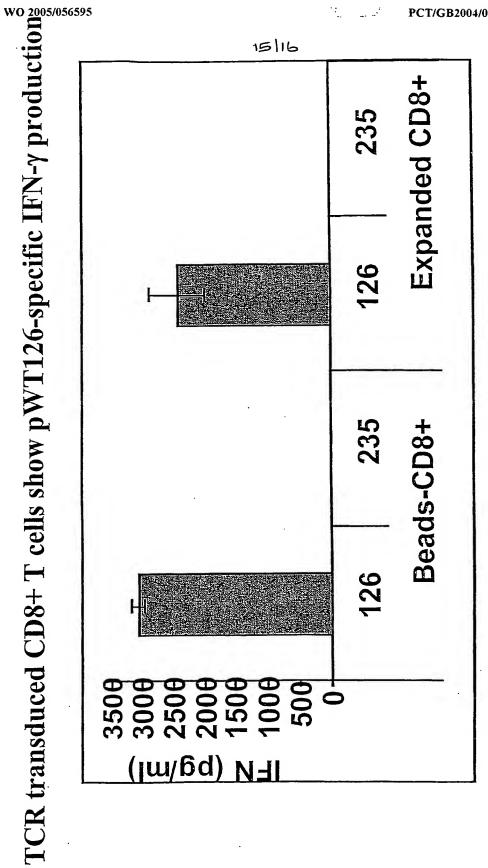
TCR transduced CD4+ T cells show pWT126-specific killing activity



% SEECHIC KILLING

e **4**

Figure



After 20 hrs incubation

16|16 Figure

